

#6

## SEQUENCE LISTING

&lt;110&gt; Mockel, Bettina

Pfefferle, Walter

&lt;120&gt; Novel Nucleotide Sequences Encoding the pfkA Gene

&lt;130&gt; 21123/274441

&lt;140&gt; 09/715,035

&lt;141&gt; 2000-11-20

&lt;160&gt; 4

&lt;170&gt; PatentIn version 3.0

&lt;210&gt; 1

&lt;211&gt; 1274

&lt;212&gt; DNA

&lt;213&gt; Corynebacterium glutamicum

&lt;220&gt;

&lt;221&gt; CDS

&lt;222&gt; (143)..(1171)

&lt;400&gt; 1

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tgatgggttt aatatggaag ac atg cga att gct act ctc acg tca ggc ggc 172

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1 5 10

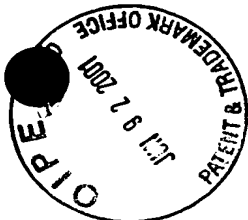
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Ser Asn Glu Phe Gly Ser Thr Val Val Gly Tyr Gln Asp Gly Trp Glu  
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09715035-062901



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cat ccg gac aag ttt aag gcc gga att gat cag att aag gcc aac tta His Pro Asp Lys Phe Lys Ala Gly Ile Asp Gln Ile Lys Ala Asn Leu 75 80 85 90	412
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ctg aag ggt gcc aag tgg ctg tct gat aac ggt atc cct gtt gtc ggt Leu Lys Gly Ala Lys Trp Leu Ser Asp Asn Gly Ile Pro Val Val Gly 110 115 120	508
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 Arg Thr Thr Val Leu Gly His Ile Gln Arg Gly Gly Thr Pro Thr Ala  
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ttc gac cgt gtt ctg gcc act cgt tat ggt gtt cgt gca gct cgt gcg 1036  
 Phe Asp Arg Val Leu Ala Thr Arg Tyr Gly Val Arg Ala Ala Arg Ala  
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tgc cat gag gga agc ttt gac aag gtt gtt gct ttg aag ggt gag agc 1084  
 Cys His Glu Gly Ser Phe Asp Lys Val Val Ala Leu Lys Gly Glu Ser  
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<212> PRT

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 35 40 45

Arg Val Gln Leu Tyr Asp Asp Glu Asp Ile Asp Arg Ile Leu Leu Arg  
 50 55 60

Gly Gly Thr Ile Leu Gly Thr Gly Arg Leu His Pro Asp Lys Phe Lys  
 65 70 75 80

062290-5E051260

Al  
ant

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Ala Leu Ile Pro Ile Gly Gly Glu Gly Thr Leu Lys Gly Ala Lys Trp  
100 105 110

Leu Ser Asp Asn Gly Ile Pro Val Val Gly Val Pro Lys Thr Ile Asp  
115 120 125

Asn Asp Val Asn Gly Thr Asp Phe Thr Phe Gly Phe Asp Thr Ala Val  
130 135 140

Ala Val Ala Thr Asp Ala Val Asp Arg Leu His Thr Thr Ala Glu Ser  
145 150 155 160

His Asn Arg Val Met Ile Val Glu Val Met Gly Arg His Val Gly Trp  
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Ile Ala Leu His Ala Gly Met Ala Gly Gly Ala His Tyr Thr Val Ile  
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Pro Glu Val Pro Phe Asp Ile Ala Glu Ile Cys Lys Ala Met Glu Arg  
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Ala Leu Pro Arg Glu Gly Thr Met Glu Leu Arg Glu Gly His Ile Asp  
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Gln Phe Gly His Lys Thr Phe Thr Gly Ile Gly Gln Gln Ile Ala Asp  
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Glu Ile His Val Arg Leu Gly His Asp Val Arg Thr Thr Val Leu Gly  
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His Ile Gln Arg Gly Gly Thr Pro Thr Ala Phe Asp Arg Val Leu Ala  
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Thr Arg Tyr Gly Val Arg Ala Ala Arg Ala Cys His Glu Gly Ser Phe  
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A!  
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<210> 4

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09715035.062500

al  
ant